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**Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease**

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2287	Tau overexpression impacts a neuroinflammation gene expression network perturbed in Alzheimer's disease. <b>2014</b> , 9, e106050		76
2286	Association between NME8 locus polymorphism and cognitive decline, cerebrospinal fluid and neuroimaging biomarkers in Alzheimer's disease. <b>2014</b> , 9, e114777		31
2285	Clinical genetics of Alzheimer's disease. <b>2014</b> , 2014, 291862		31
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2283	The impact of the human genome project on complex disease. <b>2014</b> , 5, 518-35		12
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1465	Late onset Alzheimer's disease genetics implicates microglial pathways in disease risk. <b>2017</b> , 12, 43	242
1464	Genetic variation in neurodegenerative diseases and its accessibility in the model organism <i>Caenorhabditis elegans</i> . <b>2017</b> , 11, 12	13
1463	Association of anti-inflammatory cytokine IL10 polymorphisms with motoric cognitive risk syndrome in an Ashkenazi Jewish population. <b>2017</b> , 58, 238.e1-238.e8	9
1462	Network approaches to systems biology analysis of complex disease: integrative methods for multi-omics data. <b>2018</b> , 19, 1370-1381	106

1461	Cross-Phenotype Polygenic Risk Score Analysis of Persistent Post-Concussive Symptoms in U.S. Army Soldiers with Deployment-Acquired Traumatic Brain Injury. <b>2017</b> , 34, 781-789	20
1460	Hippocampal T cell infiltration promotes neuroinflammation and cognitive decline in a mouse model of tauopathy. <b>2017</b> , 140, 184-200	112
1459	Sorting receptor SORLA: cellular mechanisms and implications for disease. <b>2017</b> , 74, 1475-1483	25
1458	Major review: Exfoliation syndrome; advances in disease genetics, molecular biology, and epidemiology. <b>2017</b> , 154, 88-103	77
1457	Dissecting the genetics of complex traits using summary association statistics. <b>2017</b> , 18, 117-127	252
1456	Two novel loci, COBL and SLC10A2, for Alzheimer's disease in African Americans. <b>2017</b> , 13, 119-129	48
1455	Integrated late onset Alzheimer's disease (LOAD) susceptibility genes: Cholesterol metabolism and trafficking perspectives. <b>2017</b> , 597, 10-16	34
1454	Genetic variants associated with risk of Alzheimer's disease contribute to cognitive change in midlife: The Atherosclerosis Risk in Communities Study. <b>2017</b> , 174, 269-282	13
1453	Gene- gene interaction between CYP2J2 and PPAR- $\alpha$ gene on late-onset Alzheimer's disease in the eastern Chinese Han population. <b>2017</b> , 322, 362-367	10
1452	Rare Genetic Variants of the Transthyretin Gene Are Associated with Alzheimer's Disease in Han Chinese. <b>2017</b> , 54, 5192-5200	16
1451	Changes in cognitive functions and cerebral grey matter and their associations with inflammatory markers, endocrine markers, and APOE genotypes in testicular cancer patients undergoing treatment. <b>2017</b> , 11, 769-783	46
1450	Multi-modal classification of Alzheimer's disease using nonlinear graph fusion. <b>2017</b> , 63, 171-181	101
1449	Polygenic score prediction captures nearly all common genetic risk for Alzheimer's disease. <b>2017</b> , 49, 214.e7-214.e11	101
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1447	Genetic variation associated with the occurrence and progression of neurological disorders. <b>2017</b> , 61, 243-264	14
1446	LD Hub: a centralized database and web interface to perform LD score regression that maximizes the potential of summary level GWAS data for SNP heritability and genetic correlation analysis. <b>2017</b> , 33, 272-279	541
1445	The ACE Gene Is Associated with Late-Life Major Depression and Age at Dementia Onset in a Population-Based Cohort. <b>2017</b> , 25, 170-177	9
1444	Exome-sequencing in a large population-based study reveals a rare Asn396Ser variant in the LIPG gene associated with depressive symptoms. <b>2017</b> , 22, 537-543	35



1443	Dementia-related adverse events in PARADIGM-HF and other trials in heart failure with reduced ejection fraction. <b>2017</b> , 19, 129-137	68
1442	DNA methylation alterations in Alzheimer's disease. <b>2017</b> , 3, dx008	26
1441	Genetic Stratification to Identify Risk Groups for Alzheimer's Disease. <b>2017</b> , 57, 275-283	21
1440	Microglia-derived ASC specks cross-seed amyloid- $\beta$ in Alzheimer's disease. <b>2017</b> , 552, 355-361	411
1439	A human microglia-like cellular model for assessing the effects of neurodegenerative disease gene variants. <b>2017</b> , 9,	59
1438	47th Annual Meeting of the German Society for Immunology, 12-15 September, 2017, Erlangen, Germany. <b>2017</b> , 47 Suppl 2, 1-333	1
1437	Comparative profiling of cortical gene expression in Alzheimer's disease patients and mouse models demonstrates a link between amyloidosis and neuroinflammation. <b>2017</b> , 7, 17762	67
1436	Genetic approaches to neurodegenerative disease. 57-76	
1435	Alzheimer's disease. 123-142	
1434	Implication of the APP Gene in Intellectual Abilities. <b>2017</b> , 59, 723-735	8
1433	Analytical Strategy to Prioritize Alzheimer's Disease Candidate Genes in Gene Regulatory Networks Using Public Expression Data. <b>2017</b> , 59, 1237-1254	12
1432	Systems healthcare: a holistic paradigm for tomorrow. <b>2017</b> , 11, 142	16
1431	Low LDL cholesterol, and genetic variation, and risk of Alzheimer's disease and Parkinson's disease: Mendelian randomisation study. <b>2017</b> , 357, j1648	90
1430	Association and interaction effects of Alzheimer's disease-associated genes and lifestyle on cognitive aging in older adults in a Taiwanese population. <b>2017</b> , 8, 24077-24087	33
1429	Aging and Neurological Diseases. <b>2017</b> ,	7
1428	Genetic Biomarkers on Age-Related Cognitive Decline. <b>2017</b> , 8, 247	23
1427	The Neurogenetics of Parkinson's Disease and Putative Links to Other Neurodegenerative Disorders. <b>2017</b> , 1-40	1
1426	Driver or Passenger: Epigenomes in Alzheimer's Disease. <b>2017</b> , 1, 5	2

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1407	Identification of genes associated with dissociation of cognitive performance and neuropathological burden: Multistep analysis of genetic, epigenetic, and transcriptional data. <b>2017</b> , 14, e1002287	56
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1390	Investigating the impact human protein-protein interaction networks have on disease-gene analysis. <b>2018</b> , 9, 455-464	7

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1388	Alzheimer risk loci and associated neuropathology in a population-based study (Vantaa 85+). <b>2018</b> , 4, e211	13
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1366	Genetic instrumental variable regression: Explaining socioeconomic and health outcomes in nonexperimental data. <b>2018</b> , 115, E4970-E4979	38
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1337	The Wisconsin Registry for Alzheimer's Prevention: A review of findings and current directions. <b>2018</b> , 10, 130-142	97
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1331	Cracking novel shared targets between epilepsy and Alzheimer's disease: need of the hour. <b>2018</b> , 29, 425-442	14
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1325	Isoprenoids and protein prenylation: implications in the pathogenesis and therapeutic intervention of Alzheimer's disease. <b>2018</b> , 53, 279-310	54
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1319	Microangiopathie cérébrale et démence : de la physiopathologie aux perspectives thérapeutiques. <b>2018</b> , 9, 140-144	
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944	Human iPSC application in Alzheimer's disease and Tau-related neurodegenerative diseases. <b>2019</b> , 699, 31-40	17
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934	Spinning convincing stories for both true and false association signals. <b>2019</b> , 43, 356-364	5
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865	A Quarter Century of APOE and Alzheimer's Disease: Progress to Date and the Path Forward. <b>2019</b> , 101, 820-838	160
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858	Blood-brain barrier transcytosis genes, risk of dementia and stroke: a prospective cohort study of 74,754 individuals. <b>2019</b> , 34, 579-590	19
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